6,500

### STIC Biotechnology Systems Branch

# RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/551.874
Source:	Pullo
Date Processed by STIC:	10/17/05
	<del></del>

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.2.2 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

### Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/55/,874
ATTN: NEW RULES CASES	S: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
	The number/text at the end of each line "wrapped" down to the next line. This may occur if your tile was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5 <sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers: use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 , "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:  (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  This sequence is intentionally skipped
8 Skipped Sequences (NEW RULES)	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.  Sequence(s) 69 missing. If intentional, please insert the following lines for each skipped sequence.  <10> sequence id number <400> sequence id number 000
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing.  Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
11Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
oug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13 Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid

AMC - Biotechnology Systems Branch - 09/09/2003



RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/551,874

DATE: 10/17/2005

TIME: 12:14:30

Input Set : N:\FANTU\10551874.raw.txt Output Set: N:\CRF4\10172005\J551874.raw use English for a U.S.

3 <110> APPLICANT: SIRS-Lab, GmbH 5 <120 > TITLE OF INVENTION: Verfahren zur Erkennung akuter generalisierter entzundlicher Zustande (SIRS), Sepsis, sepsisahnlichen Zustanden und systemischen Infektionen

8 <130> FILE REFERENCE: SL0505 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/551,874

C--> 11 <141> CURRENT FILING DATE: 2005-09-30

13 <160> NUMBER OF SEQ ID NOS: 10540

15 <170> SOFTWARE: PatentIn version 3.1

#### ERRORED SEQUENCES

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/551,874

DATE: 10/17/2005 TIME: 12:14:42

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Output Set: N:\CRF4\10172005\J551874.raw

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/551,874

DATE: 10/17/2005

TIME: 12:14:47

see p. 20 ( next page)

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Use of n and/or Xaa has been detected in the Sequence Listing Review the Sequence Listing to insure a corresponding

explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

Use of a and/or Xua has been detected in the Sequence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

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10/551,874 20

### SEQUENCE LISTING

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<120> Verfahren zur Erkennung akuter generalisierter entzündlicher Zustände (SIRS), Sepsis, sepsisähnlichen Zuständen und systemischen Infektionen

<130> SL0505

1507 (140) PCT/EP2004/0034190

577 (<141>) 2004-03-31

## **VERIFICATION SUMMARY**PATENT APPLICATION: **US/10/551,874**DATE: 10/17/2005 TIME: 12:14:48

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L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:120
L:280 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:480
M:341 Repeated in SeqNo=10
L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:420
L:507 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:60
L:683 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:180
M:341 Repeated in SeqNo=24
L:800 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:360
L:1109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:360
L:1484 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:120
M:341 Repeated in SeqNo=42
L:1516 M:341 W: (46) "n" or "Xaa" used, for SEQ ID\#:43 after pos.:180
L:1711 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:300
M:341 Repeated in SeqNo=50
L:1846 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:420
M:341 Repeated in SeqNo=55
L:1914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57 after pos.:180
M:341 Repeated in SeqNo=57
L:2069 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63 after pos.:180
L:2157 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:420
L:2193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:480
L:2224 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:240
M:341 Repeated in SeqNo=68
L\!:\!2266 M\!:\!341 W\!: (46) "n" or "Xaa" used, for SEQ ID#:69 after pos.:480
L:2292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 after pos.:240
L:2339 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71 after pos.:360
L:2531 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 after pos.:360
L:2675 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:420
L:2722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85 after pos.:0
L:2754 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:0
L:3062 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:97 after pos.:480
L:3080 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 after pos.:0
L\!:\!3147 M\!:\!341 W\!: (46) "n" or "Xaa" used, for SEQ ID#:99 after pos.:240
M:341 Repeated in SeqNo=99
L:3194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:100 after pos.:0
M:341 Repeated in SeqNo=100
L:3273 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:101 after pos.:540
M:341 Repeated in SeqNo=101
L:3589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:110 after pos.:480
M:341 Repeated in SeqNo=110
L:3689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:113 after pos.:480
M:341 Repeated in SeqNo=113
L:3778 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:116 after pos.:540
M:341 Repeated in SeqNo=116
L:4177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:118 after pos.:0
M:341 Repeated in SeqNo=118
```

#### VERIFICATION SUMMARY

PATENT APPLICATION: US/10/551,874

DATE: 10/17/2005 TIME: 12:14:48

Input Set : N:\FANTU\10551874.raw.txt
Output Set: N:\CRF4\10172005\J551874.raw

L:4273 M:341 W: (46) "n" or "Xaa" used, for SEO ID#:121 after pos.:420 L:4347 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:123 after pos.:600 M:341 Repeated in SeqNo=123 L:4575 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:132 after pos.:420 L:260276 M:301 E: (44) No Sequence Data was Shown, SEQ ID:6691 L:262782 M:301 E: (44) No Sequence Data was Shown, SEQ ID:6746 L:267220 M:301 E: (44) No Sequence Data was Shown, SEQ ID:6841 L:291623 M:301 E: (44) No Sequence Data was Shown, SEQ ID:7368 L:291985 M:301 E: (44) No Sequence Data was Shown, SEQ ID:7378 L:303036 M:301 E: (44) No Sequence Data was Shown, SEQ ID:7607 L:341027 M:301 E: (44) No Sequence Data was Shown, SEQ ID:8489 L:361142 M:301 E: (44) No Sequence Data was Shown, SEQ ID:9207 L:362191 M:301 E: (44) No Sequence Data was Shown, SEQ ID:9241 L:404719 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:14392 SEQ:10521 L:405228 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:14896 SEQ:10522 L:405801 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:16556 SEQ:10523